- A method of regulating endothelial cell growth, comprising the step of 1. contacting endothelial cells with a composition comprising a purified polypeptide in an amount effective to regulate endothelial cell growth, wherein said polypeptide:
- (a) binds the extracellular domain of Flt4 receptor tyrosine kinase and stimulates Flt4 autophosphorylation;
- (b) has an apparent molecular weight of about 23 kD as assessed by SDS-PAGE under reducing conditions; and
- (c) has an amino acid sequence comprising a portion of SEQ ID NO: 8 effective to permit binding to the Flt4 extracellular domain.
- A method according to claim 1, wherein amino terminal amino acids 2 2. through 18 of said polypeptide have an amino acid sequence corresponding to amino acids 2 through 18 set forth in SEQ ID NO: 5.
- A method according to claim 1, wherein said polypeptide is purifyable 3. from conditioned media from a PC-3 prostatic adenocarcinoma cell line, said cell line having ATCC CRL No. 1435, using an affinity purification procedure wherein the affinity purification matrix comprises a polypeptide comprising the extracellular domain of Flt4 receptor tyrosine kinase.
- A method according to claim 1 wherein the endothelial cells are lymphatic 4. endothelial cells.
- A method of modulating the activity of Flt4 receptor tyrosine kinase (Flt4), 5. comprising the steps of:

identifying a patient in need of modulation of Flt4 activity; and

administering to the patient a composition comprising a purified polypeptide in an amount effective to modulate the activity of Flt4, wherein the polypeptide is selected from the group consisting of:

(a) a polypeptide that binds the extracellular domain (EC) of Flt4, said polypeptide comprising an amino acid sequence comprising a portion of SEQ ID NO: 8 effective to permit such binding; and

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(b) an antibody which is specifically reactive with the polypeptide of (a).

A method apporting to claim 5, wherein the composition comprises a

6. A method according to claim 5, wherein the composition comprises a polypeptide that binds the extracellular domain of Flt4, said polypeptide comprising a portion of SEQ ID NO: 8 effective to permit such binding.

7. A method according to claim 5, wherein the composition further comprises a pharmaceutically-acceptable diluent, adjuvant, or carrier.

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- 8. A method according to claim 5, wherein the identifying step comprises identifying a patient suffering from a disorder of the lymphatic system, and wherein the polypeptide is administered in an amount effective to modulate Flt4 activity in endothelial cells of lymphatic vessels of the patient.
- 9. A method according to claim 5, wherein the polypeptide binds Flt4 and promotes proliferation of lymphatic endothelial cells that express Flt4.
- 10. A method according to claim 5, wherein the polypeptide binds the extracellular domain of Flt4 and stimulates Flt4 phosphorylation in mammalian cells expressing Flt4.
- 11. A method according to claim 5, wherein the polypeptide comprises a contiguous portion of SEQ ID NO: 8 that is sufficient to bind human Flt4EC,

wherein said contiguous portion includes eight cysteine residues that are conserved in human vascular endothelial growth factor (VEGF), human platelet derived growth factor A (PDGF-A), and human platelet derived growth factor B (PDGF-B), and wherein said polypeptide lacks any portion of SEQ ID NO: 8 that has one or more cysteine motifs of a Balbiani ring 3 protein (BR3P).

12. A method according to claim 5, wherein the polypeptide comprises a portion of the amino acid sequence in SEQ ID NO: 8 effective to permit said binding to

500037 method according to claim 5, wherein the polypeptide is purifyable from conditioned media, from a PC-3 prostatic adenocarcinoma cell line, said cell line having ATCC Accession Number CRL 1435, using an affinity purification procedure wherein the 5 affinity purification matrix comprises a polypeptide comprising the extracellular domain of Flt4 receptor tyrosine kinase.

- A method according to claim 5, wherein the polypeptide has an amino acid 14. sequence consisting of a portion of the amino acid sequence set forth in SEQ ID NO: 8, said portion including from residue 161 of SEQ ID NO: 8 to residue 211 of SEQ ID NO: 8, said portion lacking at least carboxy-terminal residues of SEQ ID NO: 8 beyond residue 227.
- A method according to claim 14, wherein the portion of the amino acid 15. sequence set forth in SEQ ID NO: 8 includes from residue 131 of SEQ ID NO: 8 to residue 211 of SEQ ID NO: 8.
- A method according to claim 14, wherein the portion of the amino acid 16. sequence set forth in SEQ ID NO: 8 includes from residue 113 of SEQ ID NO: 8 to residue 213 of SEQ ID NO: 8.
- A method according to claim 14, wherein the portion of the amino acid 17. sequence set forth in SEQ ID NO: 8 includes amino acids 103 to 217 of SEQ ID NO: 8.
- A method according to claim 14, wherein the portion of the amino acid 18. sequence set forth in SEQ ID NO: 8 includes amino acids 32 to 227 of SEQ ID NO: 8.
- A method of modulating the activity of Flt4 receptor tyrosine kinase (Flt4) 19. 2 wh 133/ in Flt4-expressing cells, comprising the steps of:

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-115-(a) preparing a polynucleotide comprising a nucleotide sequence that encodes a polypeptide that binds to the extracellular domain of human Flt4, wherein said polynucleotide includes a strand that hybridizes to a DNA comprising the non-coding strand complementary to SEQ ID NO: 32, under the following hybridization conditions: 5 (i) hybridization at 42 °C for 20 hours in a solution containing 50% formamide, 5x SSPE, 5x Denhardt's solution, 0.1% SDS and 0.1 mg/ml denatured salmon sperm DNA; and (ii) washing the filter twice for thirty minutes at room temperature and twice for thirty minutes at $65\,^{\circ}\text{C}$ with a wash solution containing 1x10 SSC, and 0.1% SDS; (b) transforming or transfecting a cell with the polynucleotide such that the cell expresses and secretes a polypeptide encoded by said polynucleotide, wherein said secreted polypeptide binds the extracellular domain of human Flt4 and has a molecular weight of about 23 kD as assessed by SDS-PAGE under reducing conditions; and 15 (c) contacting Flt4-expressing cells with the secreted 23 kD polypeptide. A method according to claim 19, wherein the polypeptide has an amino 20. acid sequence comprising a continuous portion of the amino acid sequence shown in SEQ ID NO: 8 effective to permit said binding. A method according to claim 19, wherein the polynucleotide comprises a 21. 20 nucleotide sequence that encodes the amino acid sequence shown in SEQ ID NO: 8, wherein the polynucleotide is transcribed and translated in the cell to produce a prepro-VEGF-C polypeptide having the amino acid sequence shown in SEQ ID No: 8, and wherein the prepro-VEGF-C polypeptide is proteolytically processed to form the 23 kD polypeptide. 25 A method according to claim 19, wherein the polynucleotide comprises the 22. polypeptide-encoding insert of plasmid pFLT4-L, deposited as ATCC Accession No. 97231.

-116-A method according to claim 19, wherein the polynucleotide further 23. includes an expression control sequence operably linked to the sequence that encodes the polypeptide. A method according to claim 19, wherein the transforming step comprises 24. contacting the cell with vector that contains the polynucleotide. 5 A method according to claim 19, wherein the Flt4-expressing cells are 25. human endothelial cells. A method according to claim 25, wherein the human endothelial cells are 26. lymphatic endothelial cells. A method according to claim 25, wherein steps (b) and (c) are performed 27. 10 in vivo. A method of modulating the proliferation of mammalian endothelial cells 28. comprising the step of contacting mammalian endothelial cells with a composition comprising a polypeptide in an amount effective to modulate the proliferation of mammalian endothelial cells, said polypeptide comprising a VEGF-C ΔC_{156} polypeptide 15 that binds to human Flt4 receptor tyrosine kinase (Flt4) and fails to bind to human KDR receptor tyrosine (VEGFR-2), said polypeptide having an amino acid sequence comprising a portion of \$EQ ID NO: 8 effective to permit binding to Flt4, wherein the cysteine residue at position 156 of SEQ ID NO: 8 has been deleted or replaced by another amino acid. 20 A method according to claim 28, wherein the portion of SEQ ID NO: 8 is 29. selected from the group consisting of: a continuous portion having as its amino terminal residue an amino acid (a) between residues 102 and 114 of SEQ ID NO: 8 and having as its carboxy terminal residue an amino acid between residues 212 and 228 of SEQ ID NO: 8, wherein the 25

cysteine residue at position 156 of SEQ ID NO: 8 has been deleted or replaced by another amino acid;

- continuous portions that comprise an amino-terminal truncation of (a); and (b)
- continuous portions that comprise a carboxyl-terminal truncation of (a) or (c) (b).
 - A method according to claim 28, wherein said endothelial cells are 30. lymphatic endothelial cells.
- An in vivo method according to claim 28, wherein the contacting step 31. comprises administering to a mammalian subject in need of modulation of the growth of lymphatic endothelial cells a composition comprising said polypeptide, in an amount 10 effective to modulate the growth of lymphatic endothelial cells in vivo.
 - A method according to claim 31, wherein said polypeptide has reduced 32. effect on the permeability of mammalian blood vessels compared to a wildtype VEGF-C polypeptide with an amino acid sequence set forth in SEQ ID NO: 8 from residue 103 to residue 227.

A method of modulating the proliferation of mammalian endothelial cells 33. comprising the step of contacting mammalian endothelial cells with a composition comprising a polypeptide in an amount effective to modulate the proliferation of mammalian endothelial cells, said polypeptide comprising a fragment of a vertebrate prepro-VEGF-amino acid sequence that binds to human Flt4 receptor tyrosine kinase, with the proviso that, in said polypeptide, a conserved cysteine of the vertebrate prepro-VEGF-C has been deleted or replaced by another amino acid,

wherein the vertebrate preprd-VEGF-C amino acid sequence comprises an amino acid sequence that is encoded by a DNA of vertebrate origin which hybridizes to a

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non-coding strand complementary to nucleotides 352 to 1611 of SEQ ID NO: 7 under the following hybridization conditions: hybridization at 42 C in a hybridization solution comprising 50% formamide, 5 X SSC, 20 mM Na•PO₄, pH 6.8; and washing in 0.2 X SSC at 55°C,

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wherein nucleotides 352 to 1611 of SEQ ID NO: 7 encode a human prepro-VEGF-C having the amino acid sequence set forth in SEQ ID NO: 8 that is characterized by eight cysteine residues at positions 131, 156, 162, 165, 166, 173, 209, and 211 of SEQ ID NO: 8 that are conserved in human vascular endothelial growth factor (VEGF), human platelet derived growth factors A and B (PDGF-A, PDGF-B), human placenta growth factor (PIGF-1), and human vascular endothelial growth factor B (VEGF-B), and

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wherein the conserved cysteine that has been deleted or replaced corresponds to position 156 of SEQ ID NO: 8.

- A method according to claim 33, wherein the vertebrate is a human. 34.
- A method according to claim 33, wherein the vertebrate is a mouse. 35.

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- A method according to claim 33, wherein said polypeptide comprises an 36. amino acid sequence selected from the group consisting of:
- the amino acid sequence of SEQ ID NO: 8, wherein the cysteine residue at (a) position 156 of SEQ ID NO: 8 has been deleted or replaced by another amino acid;
- the amino acid sequence of SEQ ID NO: 11, wherein the cysteine residue (b) at position 152 of SEQ ID NO: 11 has been deleted or replaced by another amino acid;
- the amino acid sequence of SEQ ID NO: 13, wherein the cysteine residue (c) at position 155 of SEQ ID NO: 13 has been deleted or replaced by another amino acid;
 - amino-terminal truncations of (a), (b), or (c); and (d)
 - carboxyl-terminal truncations of (a), (b), (c), or (d). (e)

An in vivo method according to claim 33, wherein the contacting step comprises administering to a mammalian subject in need of modulation of the growth of lymphatic endothelial cells a composition comprising said polypeptide, in an amount effective to modulate the growth of lymphatic endothelial cells in vivo.

38. A method for screening for inhibitors of the Flt4 receptor tyrosine kinase (Flt4), comprising the steps of:

contacting a cell that expresses Flt4 with a Flt4 ligand in the presence and absence of a putative inhibitor compound; and

assaying the Flt4 for autophosphorylation, wherein reduced autophosphorylation in the presence of the putative inhibitor compound versus the absence is identified as Flt4 inhibitory activity.

- 39. A method according to claim 38, wherein said Flt4 ligand is a polypeptide that:
- (a) binds the extracellular domain of Flt4 receptor tyrosine kinase and stimulates Flt4 autophosphorylation;
- (b) has an apparent molecular weight of about 23 kD as assessed by SDS-PAGE under reducing conditions; and
- (c) has an amino acid sequence comprising a portion of SEQ ID NO: 8 effective to permit binding to the Flt4 extracellular domain.

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